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Title

Author Correction: Gut microbiome composition in the Hispanic Community Health Study/Study of Latinos is shaped by geographic relocation, environmental factors, and obesity.

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Kaplan, Robert C
Wang, Zheng
Usyk, Mykhaylo
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AUTHOR CORRECTION

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Author Correction: Gut microbiome composition in the Hispanic Community Health Study/Study of Latinos is shaped by geographic relocation, environmental factors, and obesity

Robert C. Kaplan^{1,2*}, Zheng Wang¹, Mykhaylo Usyk³, Daniela Sotres-Alvarez⁴, Martha L. Daviglius⁵, Neil Schneiderman⁶, Gregory A. Talavera⁷, Marc D. Gellman⁶, Bharat Thyagarajan⁸, Jee-Young Moon¹, Yoshiki Vázquez-Baeza^{9,10}, Daniel McDonald¹¹, Jessica S. Williams-Nguyen², Michael C. Wu², Kari E. North¹², Justin Shaffer¹¹, Christopher C. Sollecito³, Qibin Qi¹, Carmen R. Isasi¹, Tao Wang¹, Rob Knight^{10,11,13,14} and Robert D. Burk^{1,3,15}

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Following publication of the original paper [1], an error was reported in the third paragraph in the section “**Analysis of GMB composition and its correlates**” (page 3 of the PDF). The first sentence of the text should refer to Table 2, but mistakenly refers to Table 1.

The sentence should read:

“Fungal (ITS1) GMB populations were dominated by *Aspergillus proliferans* and *Saccharomyces cerevisiae* in both mainland US-born and Latin American-born groups (Table 2).”

As a result Table 2 appears further in the article, after the second reference to Table 2.

Author details

¹Department of Epidemiology and Population Health, Albert Einstein College of Medicine, 1300 Morris Park Avenue, Bronx, NY 10461, USA. ²Division of

Public Health Sciences, Fred Hutchinson Cancer Research Center, Seattle, WA, USA. ³Department of Pediatrics, Albert Einstein College of Medicine, Bronx, NY, USA. ⁴Department of Biostatistics, Gillings School of Global Public Health, University of North Carolina, Chapel Hill, NC, USA. ⁵Institute for Minority Health Research, University of Illinois at Chicago College of Medicine, Chicago, IL, USA. ⁶Department of Psychology, University of Miami, Miami, FL, USA. ⁷Division of Health Promotion and Behavioral Science, San Diego State University, San Diego, CA, USA. ⁸Division of Molecular Pathology and Genomics, University of Minnesota, Minneapolis, MN, USA. ⁹Jacobs School of Engineering, University of California, San Diego, La Jolla, CA, USA. ¹⁰Center for Microbiome Innovation, University of California, San Diego, La Jolla, CA, USA. ¹¹Department of Pediatrics, University of California, San Diego, La Jolla, CA, USA. ¹²Department of Epidemiology, Gillings School of Global Public Health, University of North Carolina, Chapel Hill, NC, USA. ¹³Department of Computer Science and Engineering, University of California, San Diego, La Jolla, CA, USA. ¹⁴Department of Bioengineering, University of California, San Diego, La Jolla, CA, USA. ¹⁵Department of Microbiology & Immunology, Albert Einstein College of Medicine, Bronx, NY, USA.

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1. Kaplan RC, Wang Z, Usyk M, et al. Gut microbiome composition in the Hispanic Community Health Study/Study of Latinos is shaped by geographic relocation, environmental factors, and obesity. *Genome Biol.* 2019;20:219 <https://doi.org/10.1186/s13059-019-1831-z>.

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* Correspondence: robert.kaplan@einstein.yu.edu

¹Department of Epidemiology and Population Health, Albert Einstein College of Medicine, 1300 Morris Park Avenue, Bronx, NY 10461, USA

²Division of Public Health Sciences, Fred Hutchinson Cancer Research Center, Seattle, WA, USA

Full list of author information is available at the end of the article



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